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Gen. promega clean up to clean the DNA

ReadSamples Method SaveClear Print Quit  
 Results file: A:\WORK RES Method name: A:\DEFAULT  
 Assay type: General Ratio and Concentration Units: ug/ml  
 Formula setup: VIEW Background Correction: [Yes]  
 Sampling device: One cell Concentration: [Yes]  
 Read average time: 1.50 sec Peak Pick: [No]

let's get it

Core

5K50 = 250 mg

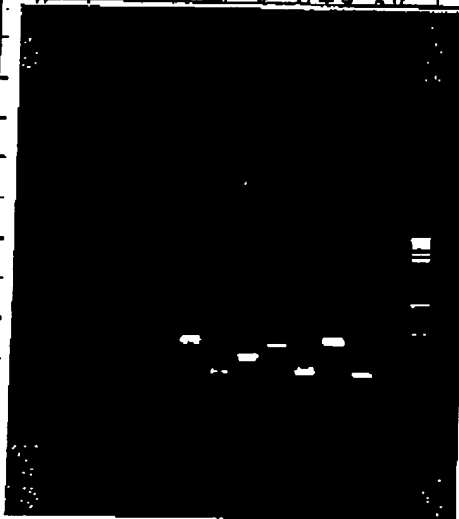
Sample ID	abs 260.0 nm	abs 280.0 nm	bkgr abs 320.0 nm	260.0 nm 280.0 nm	Protein ug/ml	Nucleic Acid ug/ml
1	-0.0114	-0.0116	-0.0120	1.2931	0.7733	0.0444
2	-0.0119	-0.0118	-0.0118	9.7100	0.1039	-0.0007
3	-0.0120	-0.0121	-0.0120	-0.0447	-22.3658	-0.0128
4	0.0478	0.0203	-0.0119	1.8583	0.5387	3.2174
5	0.0479	0.0205	-0.0119	1.8491	0.5408	3.2320
6	0.0488	0.0210	-0.0120	1.8351	0.5449	3.3033
7	0.0489	0.0210	-0.0117	1.8544	0.5393	3.2629
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Get the right sequence from DSS

28S + (4) S give recombinant product

11 10 9 8 7 6 5 4 3 2 1



1. No template negative control
2. Hygro 2 + R(-)
3. Gus + Hygro 3
4. Hygro 1 + Hygro 2
5. 28S + (4) S
6. Ac 6 + Ac 6
7. Hygro 2 + Ac (-)
8. Gus 1 + Hygro 3
9. Hygro 1 + Hygro 2
10. 28S + (4) S
11. Ac 6 + Ac 6

EXHIBIT E